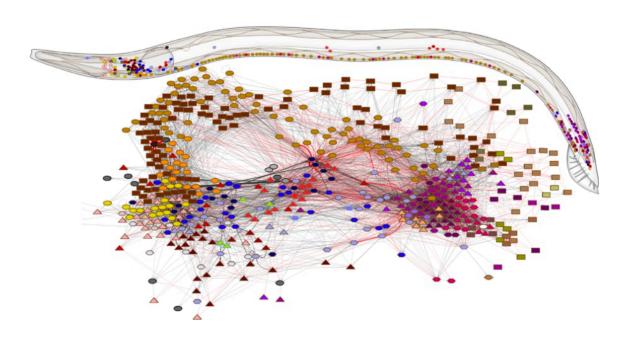


CONNECTOMICS: AURORA EARLY SCIENCE PROJECT THOMAS URAM, ALCF TURAM@ANL.GOV



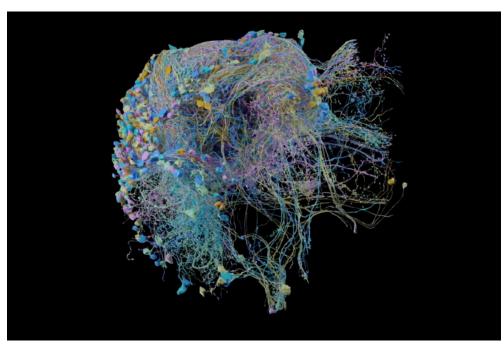
CONNECTOMICS: AURORA EARLY SCIENCE PROJECT A connectome is a comprehensive map of neural connections in the brain; in other words, its "wiring diagram".

CONNECTOMICS SO FAR



Cook, S.J., Jarrell, T.A., Brittin, C.A. et al. Whole-animal connectomes of both Caenorhabditis elegans sexes. Nature 571, 63-71 (2019)





A Connectome of the Adult Drosophila Central Brain

C. Shan Xu, Michal Januszewski, Zhiyuan Lu, Shin-ya Takemura, Kenneth

J. Hayworth, Gary Huang, Kazunori Shinomiya, Jeremy Maitin-

Shepard, David Ackerman, Stuart Berg, Tim Blakely, John Bogovic, Jody Clements, Tom Dolafi, Philip Hubbard, Dagmar Kainm ueller, William Katz, Takashi Kawase, Khaled A. Khairy, Laramie Leavitt, Peter H. Li, Larry Lindsey, Nicole Neubarth, Donald J. Olbris, Hideo Otsuna, Eric

T. Troutman, Lowell Umayam, Ting Zhao, Masayoshi Ito, Jens Goldammer, Tanya Wolff, Robert Svirskas, Philipp Schlegel, Erik a R. Neace, Christopher J. Knecht Jr., Chelsea X. Alvarado, Dennis A. Bailey, Samantha Ballinger, Jolanta A Borycz, Brandon S. Canino, Natasha Cheatham, Michael Cook, Marisa Dreher, Octave Duclos, Bryon Eubanks, Kelli Fairbanks, Samantha Finle

y, Nora Forknall, Audrey Francis, Gary Patrick Hopkins, Emily M. Joyce, SungJin Kim, Nicole A. Kirk, Julie Kovalyak, Shirley A. Lauchie, Alanna Lohff, Charli Maldonado, Emily

A. Manley, Sari McLin, Caroline Mooney, Miatta Ndama, Omotara Ogundeyi, Nneoma Okeoma, Christopher Ordish, Nicholas P adilla, Christopher Patrick, Tyler Paterson, Elliott E. Phillips, Emily M. Phillips, Neha Rampally, Caitlin Ribeiro, Madelaine K Robertson, Jon Thomson Rymer, Sean M. Ryan, Megan Sammons, Anne K. Scott, Ashley

L. Scott, Aya Shinomiya, Claire Smith, Kelsey Smith, Natalie L. Smith, Margaret

A. Sobeski, Alia Suleiman, Jackie Swift, Satoko Takemura, Iris Talebi, Dorota Tarnogorska, Emily Tenshaw, Temour Tokhi, John J. Walsh, Tansy Yang, Jane Anne Horne, Feng Li, Ruchi Parekh, Patricia

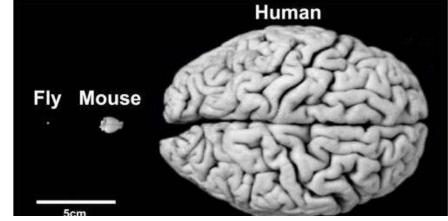
K. Rivlin, Vivek Jayaraman, Kei Ito, Stephan Saalfeld, Reed George, Ian Meinertzhagen, Gerald M. Rubin, Harald

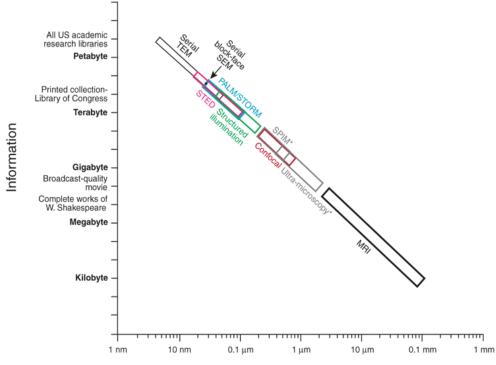
F. Hess, Louis K. Scheffer, Viren Jain, Stephen M. Plaza

bioRxiv 2020.01.21.911859; doi: https://doi.org/10.1101/2020.01.21.911859

ENABLING CONNECTOMICS AT EXASCALE TO FACILITATE DISCOVERIES IN NEUROSCIENCE

- Why Connectomics?
 - Understand neurodegenerative diseases
 - Understand learning and aging and consciousness
 - Build better, purpose-built neural networks
 - Design neuromorphic computers
- Why Exascale Connectomics?
 - Fruit fly brain: 250K neurons, O(terabytes)
 - Mouse brain: 14M neurons, O(petabytes)
 - Human brain: 100B neurons, O(exabytes)





Resolution



CONNECTOME CHALLENGES

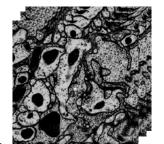
Big raw data

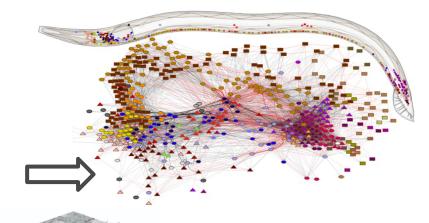
Limited training data

Significant human effort for ground truth

- Many imperfections or anomalies in data
- Complex neuron shapes in different sizes
- Deep learning model training and large-scale inference







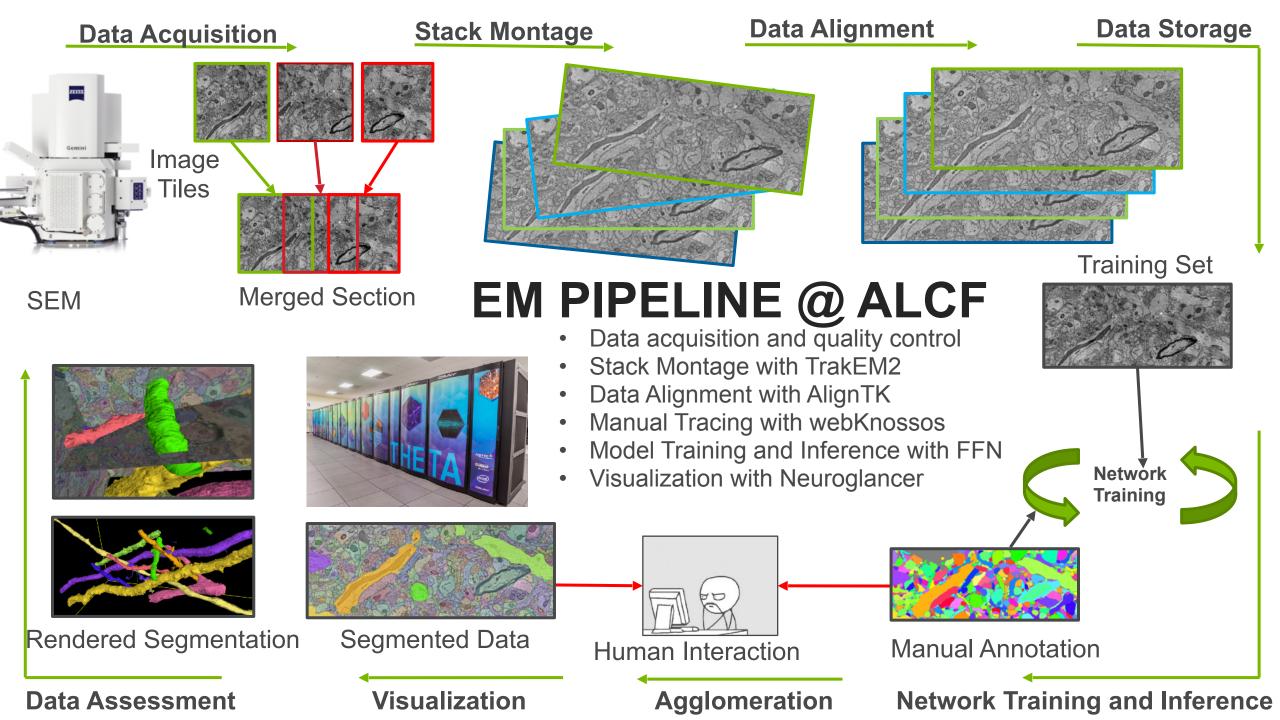




| Step | Software | Computing Env |
|--------------|----------|-----------------|
| Acquisition | ATLAS | SEM Workstation |
| Stitching | TrakEM2 | Cooley / Theta |
| Alignment | AlignTK | Theta |
| Segmentation | FFN | Theta |



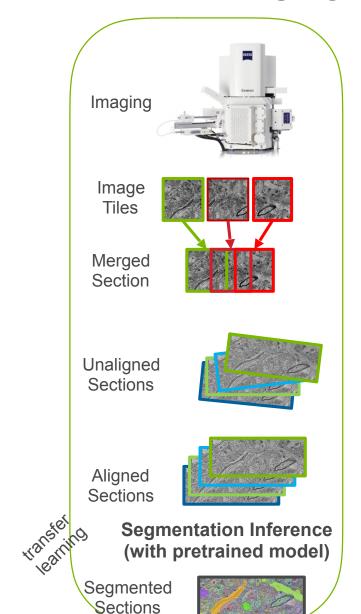


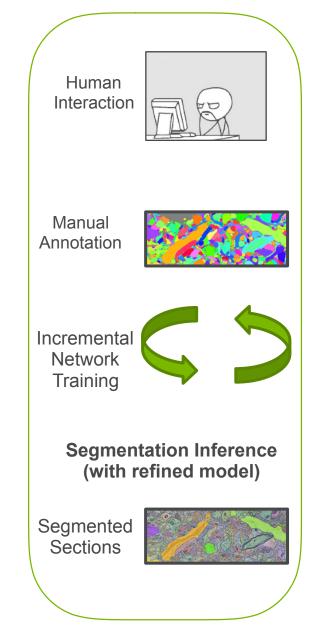


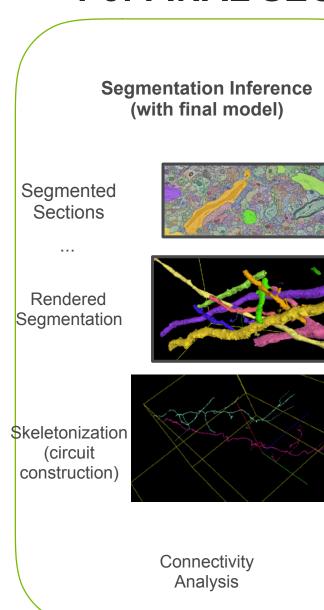
P1: INITIAL SEG

P2: REFINEMENT

P3: FINAL SEG





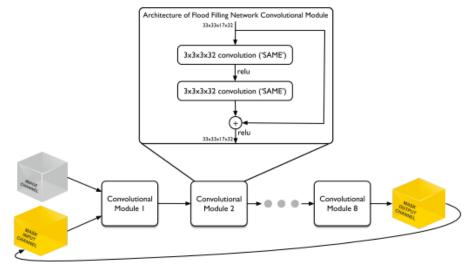


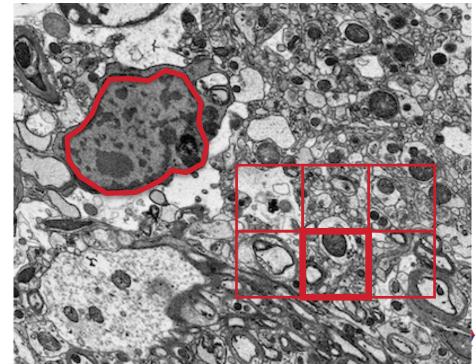
FLOOD FILLING NETWORK (FFN)

TRAINING

- FFN was developed by Google Brain for automated segmentation of structures in EM image data
- Implemented in TensorFlow
- Network consists of a series of 3D convolutional blocks with residual connections
- A 12 layer network has roughly 0.5M trainable parameters
- Network builds on notion of watershed algorithm
 - Find boundaries and fill interior
 - Prefer split errors over merge errors
- Boundary-finding is complicated
 - complex object structure and substructure
 - variation between datasets (limited opportunity for transfer learning, and yet...)
 - variation within datasets (fixing, cutting, staining, imaging)
- Dataset-specific training is required
- Accuracy of FFN is an order of magnitude better than past approaches, but with a higher computational cost.

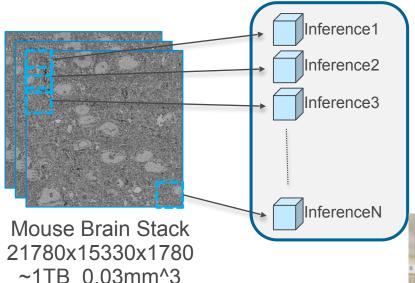
M. Januszewski, J. Kornfeld, P. H. Li, A. Pope, T. Blakely, L. Lindsey, J. Maitin-Shepard, M. Tyka, W. Denk, and V. Jain, "High-precision automated reconstruction of neurons with flood-filling networks," Nature Methods





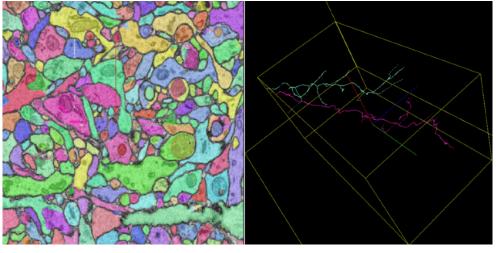
FLOOD FILLING NETWORK (FFN)

INFERENCE



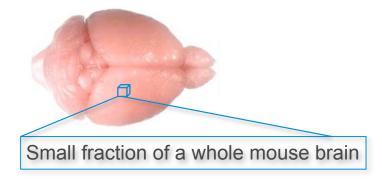
55,000 inference jobs 256^3 pixels per job 2048 nodes, 16h (including I/O)





Segmentation and Skeletonization

100x increase in computation 20x increase in I/O speed



Current processing and I/O challenges will be solved with the upgrade to Aurora.

The new system will enable the processing of whole mouse brains at high resolution.

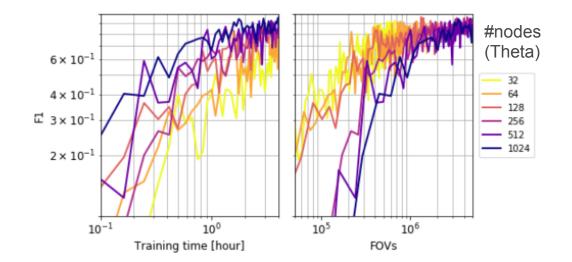


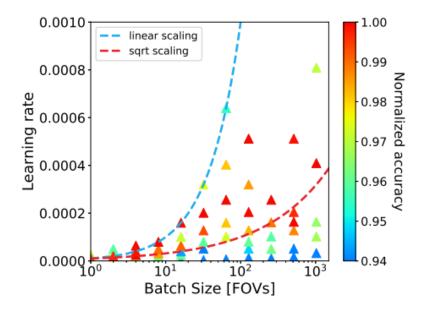


40nm resolution

ADVANCES

- Scaling FFN training and inference on ALCF machines
- Initial hyperparameter scans: batch size, learning rate, optimizer, num layers
- Hyperparameter optimization with Deep Hyper (ongoing)
- Evaluation of impact of reduced precision
- Evaluation on future architectures
- Application of learning in other parts of the pipeline (e.g. image alignment)









SUMMARY

- Significant learning and computing challenges remain
- Significant human effort still required
 - How long before learning methods outperform humans on these tasks?
- Obtaining a full mouse connectome will take years of imaging and years of computing
 - Human brain?
- Future will bring advances in microscopes (more pixels!) and methods, and advances in computing (Aurora)
- We are using brain-inspired* neural networks to understand the brain so we can make better neural networks.















































